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Conference Abstract

Extended Taxonomic Curation: Moving beyond species lists to linking species data

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Abstract

Taxonomy is at the center of modern biodiversity science. No species can be systematically studied until it is defined, and no observation can be linked to related data without a taxonomic label. However, taxonomy is also a science in constant flux—even well-studied groups like Mammalia have fluctuated by >25% in recognized species in the last decade (Burgin et al. 2018, MDD 2022a, MDD 2022b). As a result, there are calls to create a “global list of accepted species” to increase taxonomic stability, particularly for policy decisions in biodiversity conservation and management (Garnett et al. 2020). The counterargument notes that forcing definitional consensus is likely to further inequities, and that a pluralistic, coordinated approach to taxonomy can be achieved with innovative cyberinfrastructure designs and services (Sterner et al. 2020, Franz and Sterner 2018).

Here, we propose that digitally “extended” taxonomic curation can play new and innovative roles in

1. linking observational data to alternative taxonomic concepts; and
2. enabling fit-for-use taxonomy to inform policy decisions.

Taxonomic curators (TCs) have traditionally limited their activities to making lists of accepted species and higher taxa. However, most of today's biodiversity questions require observational data (e.g., specimen occurrences) that are taxonomically coherent, not just name lists, and for those linked data to be digitally available in public databases. If the

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collective activities of TCs can be effectively unified across distributed networks, they might facilitate the transition to Extended Specimen Networks of taxonomically coherent biodiversity data, a core goal of current research initiatives (e.g., Lendemer et al. (2020)).

Beyond lists of species names is the domain of what names mean in practice (i.e., taxonomic concepts), which often differs by author (Fig. 1). Here we argue that curating the various lines of evidence that represent taxonomic concepts—what we call Species Meaning Artifacts (SMArts)—is a promising strategy for keeping track of how species splits and lumps will affect observational data records in the Global Biodiversity Information Facility (GBIF) or National Center for Biotechnology Information (NCBI). Instead of labeling records by a static name, records can be digitally associated with SMArt evidence from alternative taxonomies (e.g., geographic range maps before/after a species split). Networks of TCs curating digital SMArts will enable 'taxonomically intelligent' data aggregation (Bisby 2000), a long-pursued goal in biodiversity data science that, once realized, promises to enable investigations ranging from viral spillover to biodiversity loss (Upham et al. 2021).

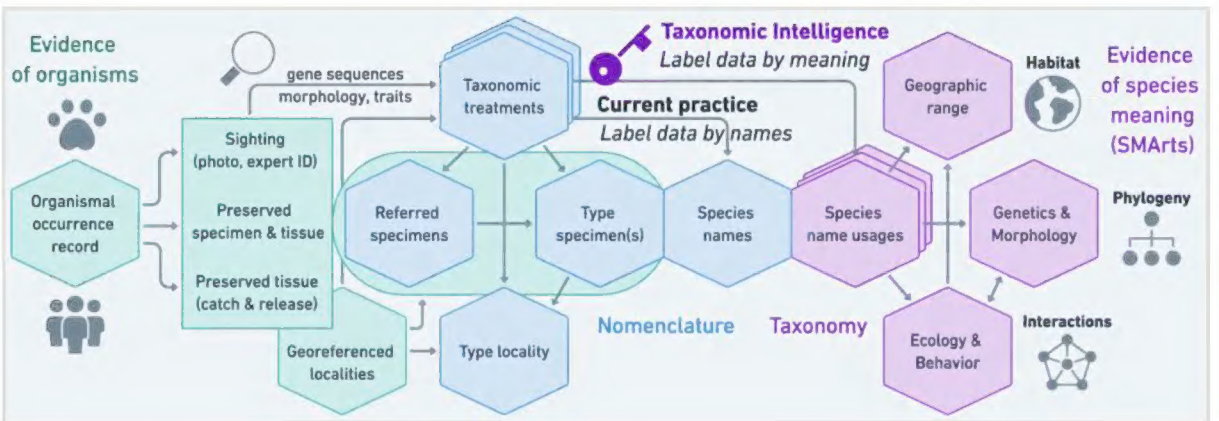

Figure 1.

Knowledge of species is rooted in organismal observations, and created via the flow of information from nomenclature (naming species) to taxonomy (defining species boundaries and relationships). Current practice is to label observational data by names alone; however, curating the lines of evidence that represent the conceptual meanings of those names, and how those meanings differ among authors through time, will allow for more accurate data labeling and aggregation (i.e., taxonomic intelligence).

Keywords

biodiversity data science, mammals, taxonomic intelligence

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